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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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301.5
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Gapop 10.0 , Gapext 0.5
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1436
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/cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09_NEW_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*
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/cgn2_6/ptodata/2/pubpaa/US00_NEW_PUB.pep:*
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     US-10-091-135-88
US-10-091-135-87
US-09-876-225-3
US-10-091-135-86
US-09-876-225-3
US-09-876-225-3
US-10-123-04-262
US-10-123-94-262
US-10-123-94-262
US-10-176-918-262
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  bpaa/US08_NEW_PUB.pep:*
Sequence 88, Appl
Sequence 2, Appl1
Sequence 3, Appl1
Sequence 3, Appl1
Sequence 33, Appl1
Sequence 262, App
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Description
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-10-031-135-88

Sequence 88, Application US/10091135

Publication No. US2003003660A1

GENERAL INFORMATION:
APPLICANT: King, Te Piao
APPLICANT: King, Te Piao
APPLICANT: Spangfort, Michael Dho
TITLE OF INVENTION: RECOMBINANT HYBRID ALLERGEN CONSTRUCTS WITH REDUCED
TITLE OF INVENTION: RECOMBINANT HYBRID ALLERGEN CONSTRUCTS WITH REDUCED
TITLE OF INVENTION: ALLERGENICITY THAT RETAIN INMUNOGENICITY OF THE NATURAL ALLE
FILE REFERENCE: 2313/1H587-US1
CURRENT APPLICATION NUMBER: US/10/091,135
CURRENT APPLICATION NUMBER: US/0/272,818
PRIOR APPLICATION NUMBER: US 60/272,818
PRIOR FILING DATE: 2001-03-02

NUMBER OF SEQ ID NOS: 98
SOFTWARE: Patentin version 3.1
SEQ ID NOS: 98
LENGTH: 222
TYPE: PRT
CORANISM: Homo sapiens
US-10-091-135-88
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Best Local Similarity
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72.5%;
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RESULT		45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30.	29	28	27	26	25	24	23	22	21	. 20
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	ALIGNMENTS	US-10-142-431-360	US-09-909-204-285	US-09-904-859-285	US-09-904-820-285	US-10-140-474-360	US-10-137-865-360	US-10-176-921-360	US-10-176-918-360	US-10-175-746-360	US-09-907-942-285	US-09-907-613-285	US-09-906-838-285	US-10-140-470-360	US-10-123-904-360	US-10-121-049-360	US-09-906-742-285	US-10-028-072-360	US-09-904-011-285	US-09-907-841-285	US-09-907-824-285	US-09-902-853-285	US-09-905-291A-285	US-10-230-338-56	US-10-218-631-56	US-10-230-163-56	US-10-227-884-56
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CURRENT APPLICATION NUMBER: US/09/876,225
CURRENT FILING DATE: 2001-06-08
PRIOR APPLICATION NUMBER: US 60/209,989
PRIOR FILING DATE: 2000-06-08
NUMBER OF SEQ ID NOS: 5
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 2
LENGTH: 255
TYPE: PRT
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US-10-091-135-87
Sequence 87, Application US/10091135
; Publication No. US20030039660A1
; GENERAL INFORMATION:
APPLICANT: King, Te Piao
APPLICANT: King, Te Piao
APPLICANT: Spangfort, Michael Dho
TITLE OF INVENTION: RECOMBINANT HYBRID ALLERGEN CONSTRUCTS WITH REDUCED
TITLE OF INVENTION: ALLERGENICITY THAT RETAIN INMUNOGENICITY OF THE NAIL
FILE REFERENCE: 2313/1H587-US1
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                                                                                                               Query Match
Best Local :
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SEQ ID NO 87
LENCTH: 223
TYPE: PRT
                                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 2, Application US/09876225 Patent No. US20020052025A1
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Matches
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TITLE OF INVENTION: RTVP Based Compositions and Methods for the Treatment of Prostate
FILE REFERENCE: 38594.0027
                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Baylor College of Medicine APPLICANT: Thompson, Timothy C. APPLICANT: Ren, Chengzhen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/091,135
CURRENT FILING DATE: 2002-03-04
PRIOR APPLICATION NUMBER: US 60/272,818
PRIOR FILING DATE: 2001-03-02
NUMBER OF SEQ ID NOS: 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            155 VVWYSSYLVGCGNAYCPNQKVLKYYYVCQYCPAGNWANRLYVPYEQGAPCASCPDNCDDG 214
                                            19 VLLFLVAGLLPSFPANEDKDPAFTALL---TTQTQVQREIVNKHNELRRAVSPPARNMLK 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 EHTSKEERTIDGVECGENLFFSSAPYTWSYAIQNWFDERKYFRFNYGPTAQNVMIGHYTQ 120
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wes 116; Conserv
                                                                                                           Local Similarity
            VILAVIVWMASSVSSS-----SFTASTLPDITNEDFIKECVQVHNQLRSKVSPPARNMLY 57
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                                                                                          Conservative
                                                                                      24.0%; Score 344; DB 10;
33.6%; Pred. No. 2.1e-24;
7ative 37; Mismatches 81;
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; Sequence 3, Application US/09876225
Patent No. US20020052025A1
GENERAL INFORMATION:
APPLICANT: Baylor College of Medicine
APPLICANT: Thompson, Timothy C.
APPLICANT: Ren, Chengzhen
TITLE OF INVENTION: RTVP Based Compositions and Methods for the Treatment of Pros
FILE REFERENCE: 38594.0027
CURRENT APPLICATION NUMBER: US/09/876,225
CURRENT FILING DATE: 2001-06-08
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TYPE: PRT
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APPLICANT: Spangfort, Michael Dho
TITLE OF INVENTION: RECOMBINANT HYBRID ALLERGEN CONSTRUCTS WITH REDUCED
TITLE OF INVENTION: ALLERGENICITY THAT RETAIN IMMUNOGENICITY OF THE NATURAL ALLE
FILE REFERENCE: 2313/14587-US1
CURRENT APPLICATION NUMBER: US/10/091,135
CURRENT FILMS DATE: 2002-09-04
PRIOR APPLICATION NUMBER: US 60/272,818
PRIOR APPLICATION NUMBER: US 60/272,818
PRIOR APPLICATION DATE: 2001-03-02
NUMBER OF SEQ 1D NOS: 98
SOFFWARE: Patentin version 3.1
SEQ ID NO 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-10-091-135-86

S-quence 86, Application US/10091135

Publication No. US20030039660A1

GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; ORGANISM: Homo sapiens US-10-091-135-86
160 YPT----WPYKRGATCSACPNNDKCLDNLCVN 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       129 WFDEYNDFDFGVGPKTPNAVVGHYTQVVWYSSYLVGCGNAVCPNQKVLKYYYVCQYCPAG 188
                                                         190 WANRLYVPYEQGAPCASCPDN--CDDGLCTN 218
                                                                                                                  104 DFKT--RICKKVCGHYTQVVWADSYKVGCAVQFCP--KVSGFDALSNGAHFICNYGPGGN 159
                                                                                                                                                                           137 DEGVGPKTPNAVVGHYTQVVWYSSYLVGCGNAYCPNQKVLKY-----YYVCQYCPAGN 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          226 FLIAKSVLLLLSVIITIWVKHK 247
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                                                                                                                                                                                                                                        46 IAKAWASNCQFSHNTRLKPPHKLHPNFTSL--GENIWTGSVPIFSVSSAITNWYDEIQDY 103
                                                                                                                                                                                                                                                                                       85 NAQKWANQCNYRHS---NPKDRM----TSLKCGENLYMSSAP-SSWSQAIQSWFDEYNDF 136
                                                                                                                                                                                                                                                                                                                                                                                                                25 AGLLPSFPANEDKDPAFTALLTTQTQVQREIVNKHNELRRAVSPPARNMLKMEWNKEAAA 84
                                                                                                                                                                                                                                                                                                                                                             1 ANILPDI-ENED------FIKDCVRIHNKFRSEVKPTASDMLYMTWDPALAQ 45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity 37.0
78; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22.8%; Score 327.5; DB 9 37.0%; Pred. No. 6.6e-23; tive 29; Mismatches 63
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B 8

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173 QKVLKY-----YYVCQYCPAGNWANRLYVPYEQGAPCASCPDN--CDDGLCTN 218

-KVSGFDALSNGAHFICNYGPGGNYPT---WPYKRGATXSACPNNDKCLDNLCVN 244

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; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-876-225-3
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                                                                                                                                                                    밁
                                                                                                                                                                                                                                                                                                                                           ; NAME/KEY: SITE ; LOCATION: (228) ; COTHER INFORMATION: Xea equals any of the naturally occurring L-amino acids US-09-925-301-853
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-925-301-853; Sequence 853; A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 853
LENGTH: 302
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Rosen et al.
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PALO6
CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT FILING DATE: 2001-08-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No. US20020052308A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local :
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                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 60/209,989
PRIOR FILING DATE: 2000-06-08
NUMBER OF SEQ ID NOS: 5
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 1694
SOFTWARE: PatentIn Ver. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   160 SGFDALSNGAHFICNYGPGGNYPT --- WPYKRGATCSACPNNDKCLDNLCVN 208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       176 LKY-----YYVCQYCPAGNWANRLYVPYEQGAPCASCPDN--CDDGLCTN 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   104 SVPIFSVSSAITNWYDEIQDYDFKT--RICKKVCGHYTQVVWADSYKVGCAVQFCP--KV 159
                        114 YMSSAP-SSWSQAIQSWFDEYNDFDFGVGFKTPNAVVGHYTQVVWYSSYLVGCGNAYCFN 172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              117 SAP-SSWSQAIQSWFDEYNDFDFGVGPKTPNAVVGHYTQVVWYSSYLVGCGNAYCPNQKV 175
137 WTGSVPIFSVSSAITNWYDEIQDYDFKT--RICKKVCGHYTQVVWADSYKVGCAVQFCP- 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          46 SEVKPTASDMLYMIWDPALAQIAKAWASNCQFSHNTRLKPPHKIHPNFTSL--GENIWTG 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   64 RAVSPPARNMLKMEWNKEAAANAQKWANQCNYRHS---NPKDRM----TSLKCGENLYMS 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 34.9%;
                                                                                    79
                                                                                                          34 ROSMRVTLATIAWMVSFVSNYSHTANILPDI-ENED------FIKDCVRIHN 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9 LETTAMTLEPVILEL-----VAGILPSEPANEDKDPAFTALLTTQTQVQREIVNKHNELR 63
                                                                                                                                                                                                                                                      Match 22.1%;
Local Similarity 35.3%;
es 83; Conservative 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 MRVTLATIAWMVSFVSNYSHTANILPDI-ENED------FIKDCVRIHNKFR 45
                                                                                                                                                                                                             2 KQILHPALETTA-KTLFPVLLFLVAGLLPSFPANEDKDPAFTALLTTQTQVQREIVNKHN 60
                                                                                  KFRSEVKPTASDKLYMTWDPALAQIAKAWASNCQFSHNTRLKPPHKLHPNFTSL--GENI 136
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                                                                                                                                                                                                                                                        31; Mismatches
                                                                                                                                                                                                                                Score 318; DB 10; Pred. No. 6.5e-22; Indels 42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 325; DB 10; Length 266;
Pred. No. 1.3e-22;
4; Mismatches 71; Indels 46;
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CURRENT EILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 05/04/9911
PRIOR APPLICATION NUMBER: 60/05/974
PRIOR APPLICATION NUMBER: 60/05/974
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PRIOR APPLICATION NUMBER: 60/05/9113
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PRIOR APPLICATION NUMBER: 60/05/9117
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PRIOR APPLICATION NUMBER: 60/05/9122
PRIOR APPLICATION NUMBER: 60/05/9184
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PRIOR APPLICATION NUMBER: 60/05/9185
PRIOR APPLICATION NUMBER: 60/05/9185
PRIOR APPLICATION NUMBER: 60/05/9352
PRIOR APPLICATION NUMBER: 60/05/9352
PRIOR APPLICATION NUMBER: 60/05/9886
PRIOR APPLICATION NUMBER: 60/05/9886
PRIOR APPLICATION NUMBER: 60/05/9886
PRIOR APPLICATION NUMBER: 60/05/286
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TITLE OF INVENTION:
FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION:
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Godowski, Paul J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Smith, Victoria
Stewart, Timothy A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gurney, Austin L.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DeForge, Laura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wood, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tumas, Daniel
Watanabe, Colin K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sherwood, Steven
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gao, Wei-Qiang
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o. US20030004311A1
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PR APPLICATION NUMBER: 60/063550

PR FILING DATE: 1997-10-28

PR APPLICATION NUMBER: 60/063561

PR FILING DATE: 1997-10-28

PR APPLICATION NUMBER: 60/063704

PR FILING DATE: 1997-10-29

PR FILING DATE: 1997-10-29

JR APPLICATION
JR FILING DATE: 1997-10-1/
DR FILING DATE: 1997-11-03
OR FILING DATE: 1997-11-03
IOR APPLICATION NUMBER: 60/064809
IOR APPLICATION NUMBER: 60/065186

FILING DATE: 1997-11-07
APPLICATION NUMBER: 60/065186
FILING DATE: 1997-11-12

APPLICATION NUMBER: FILING DATE: 1997-10 APPLICATION NUMBER:

60/063738

: 1997-10-17 NUMBER: 60/063755 1997-10APPLICATION NUMBER: 60/063733
FILING DATE: 1997-10-29
APPLICATION NUMBER: 60/063735
FILING DATE: 1997-10-29

RR APPLICATION NUMBER: 60/065846
RR FILING DATE: 1997-11-17
DR APPLICATION NUMBER: 60/066364
DR FILING DATE: 1997-11-21
RR APPLICATION NUMBER: 60/066453
DR FILING DATE: 1997-11-24

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Query Match
Best Local S
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OR PILING DATE: 1998-05-13
OR APPLICATION UNMBER: 60/08539
OR APPLICATION UNMBER: 60/08579
OR FILING DATE: 1998-05-15
OR APPLICATION UNMBER: 60/08579
OR FILING DATE: 1998-05-15
OR APPLICATION UNMBER: 60/08570
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OR APPLICATION UNMBER: 60/088810
OR APPLICATION UNMBER: 60/08959
OR APPLICATION UNMBER: 60/08959
OR APPLICATION UNMBER: 60/089947
OR APPLICATION UNMBER: 60/09044
OR APPLICATION UNMBER: 60/09049
OR FILING DATE: 1998-06-17
OR APPLICATION UNMBER: 60/09049
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RE FILING DATE: 1998-07-01
DR APPLICATION NUMBER: 60/091519
DR FILING DATE: 1998-07-02
DR APPLICATION NUMBER: 60/091982
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R FILING DATE: 1998-05-07
DR APPLICATION NUMBER: 60/08
PR FILING DATE: 1998-05-07
DR APPLICATION NUMBER: 60/08
DR FILING DATE: 1998-05-07
DR APPLICATION NUMBER: 60/08
DR FILING DATE: 1998-05-12
DR APPLICATION NUMBER: 60/08
DR FILING DATE: 1998-05-13
DR FILING DATE: 1998-05-13
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DR APPLICATION NUMBER: 60/066511

DR FILLING DATE: 1997-11-24

DR APPLICATION NUMBER: 60/066770

R FILLING DATE: 1997-12-11

DR APPLICATION NUMBER: 60/069278

DR FILLING DATE: 1997-12-11

DR APPLICATION NUMBER: 60/069278

DR APPLICATION NUMBER: 60/06934

DR FILLING DATE: 1997-12-11

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DR APPLICATION NUMBER: 60/074086

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DR APPLICATION NUMBER: 60/074092

DR FILLING DATE: 1998-02-09

DR APPLICATION NUMBER: 60/074092

DR APPLICATION NUMBER: 60/074092

DR FILLING DATE: 1998-02-09

DR APPLICATION NUMBER: 60/074092

DR FILLING DATE: 1998-02-09

DR APPLICATION NUMBER: 60/074092

DR FILLING DATE: 1998-02-09

οy 망 밁 Ş веst Local Matches 7 50 PPAADMKYMIWDKGLAKMAKAWANQCKFEHNDCLDK--SYKCYAAFEYVGENIWLGGIKS 8 N æ PPARNMLKMEWNKEAAANAQKWANQCNYRHSNPKDRMTSLKC-----GENLYMSSAPS 120 ALETTAMTLEPVILELVAGILPSEPANEDKDPAFTALLTTQTQVQREIVNKHNELRRAVS 67 ALKNKFSCLWILGLCLVATTSSKIPS -- ITDPHFI--78; Conser Conservative Score 301.5; DB 9; Pred. No. 1.7e-20; l; Mismatches 82; ------DNCIEAHNEWRGKVN 49 Indels 29; 107

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OR APPLICATION NUMBER: 60/080165
OR FILING DATE: 1998-03-31
OR APPLICATION NUMBER: 60/081203
OR FILING DATE: 1998-04-09
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OR APPLICATION NUMBER: 60/083324

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R FILING DATE: 1998-03-25
NR APPLICATION NUMBER: 60/079663
NR FILING DATE: 1998-02-27
RR FPLICATION NUMBER: 60/079728
R FPLICATION NUMBER: 60/079728
R FILING DATE: 1998-03-27

. 1998-02-27 NUMBFD

APPLICATION NUMBER: 60/077791
FILING DATE: 1998-03-12
APPLICATION NUMBER: 60/078910
FILING DATE: 1998-03-20

121 -SWSQAIQSWFDEYNDFDFGVGPKTPNAVVGHYTQVVWYSSYLVGCGNAYCPN-QKVLKY 178

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APPLICANT:

DeForge, Laura

Filvaroff, Ellen

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; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-121-049-262
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                                                                                                                  RESULT 9
US-10-123-904-262
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 550
SEQ ID NO 262
LENGTH: 242
                                             Sequence 262, Application US/10123904 Publication No. US20030022328A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C17 CURRENT APPLICATION NUMBER: US/10/121,049 CURRENT FILING DATE: 2002-04-12 Prior Application removed - See File Wrapper or Palm Number Of Colesco Common File Reports Application removed - See File Wrapper or Palm
     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Baker, Kevin P.
                                                                                                                                                                                                        166 IFVCNYGPAGNFAN--MPPYARGESCSLCSKEEKCVKNLC 203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         166 IFVCNYGPAGNFAN--MPPYARGESCSLCSKEEKCVKNLC 203
                                                                                                                                                                                                                                                      179 YYVCQYCPAGNWANRLYVPYEQGAPCASC -- PDNCDDGLC 216
                                                                                                                                                                                                                                                                                                   108 FTPRHAITAWYNETQFYDF -- DSLSCSRVCGHYTQLVWANSFYVGCAVAMCPNLGGASTA 16:
                                                                                                                                                                                                                                                                                                                                                                                            50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 ALKNKFSCLWILGLCLVATTSSKIPS--ITDPHFI-----DNCIEAHNEWRGKVN 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8 ALETTAMTLEPVILLELVAGLLPSFPANEDKDPAFTALLTTQTQVQREIVNKHNELRRAVS 67
                                                                                                                                                                                                                                                                                                                                         -SWSQAIQSWFDEYNDFDFGVGPKTPNAVVGHYTQVVWYSSYLVGCGNAYCPN-QKVLKY 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    YYVCQYCPAGNWANRLYVPYEQGAPCASC -- PDNCDDGLC 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FTPRHAITAWYNETQFYDF--DSLSCSRVCGHYTQLVWANSFYVGCAVAMCPNLGGASTA 165
                                                                                                                                                                                                                                                                                                                                                                                          PPAADMKYMIWDKGLAKMAKAWANQCKFEHNDCLDK--SYKCYAAFEYVGENIWLGGIKS 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
                         Baker, Kevin P.
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     Beresini, Maureen
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Pred. No. 1.7e-20;
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; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-123-904-262
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NUMBER OF SEQ ID NOS: 550
SEQ ID NO 262
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 262, Applic Publication No. US20 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   121 -SWSQAIQSWFDEYNDFDFGVGPKTPNAVVGHYTQVVWYSSYLVGCGNAYCPN-QKVLKY 178
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Local Similarity 35.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8 ALETTAMTLEPVLLELVAGLLPSEPANEDKDPAETALLTTQTQVQREIVNKHNELRRAVS
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INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC INVENTION: ACIDS ENCODING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                   Baker, Kevin P.
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Godowski, Paul J.
Gurney, Austin L.
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Watanabe, Colin K
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Godowski, Paul J.
Gurney, Austin L.
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                                                                                                                                         Stewart, Timothy A.
                                                                                                                                                               Smith, Victoria
                                                                                                                                                                                      Sherwood, Steven
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                                                                                             Watanabe,Colin K
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                                                                       ood, William,
                                                     Zemin
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Pred. No. 1.7e-20;
31; Mismatches 82;
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Length 242; Indels

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US-10-175-746-262
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CURRENT PILING DATE: 2002-05-06
Prior Application removed - See Palm or File Wrapper NUMBER OF SEQ ID NOS: 550
SEQ ID NO 262
SEQ ID NO 262
                                                                                                                                                                            Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 262
LENGTH: 242
TYPE: PRT
                                                           Query Match 21.0
Best Local Similarity 35.5
Matches 78; Conservative
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Best Local
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CURRENT FILING DATE: 2002-06-19
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 35.5 es 78; Conservative
REFERENCE: P3330R1C353
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -SWSQAIQSWFDEYNDFDFGVGPKTPNAVVGHYTQVVWYSSYLVGCGNAYCPN-QKVLKY 178
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                                                                                                                                                                                                                                                                                                                                                                                                                        Tumas, Daniel
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Smith, Victoria
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Godowski,Paul J.
Gurney,Austin L.
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                                                                                                                                                                                                                                                                                                                                                                                                     tanabe, Colin K
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Pred. No. 1.7e-20;
NI: Mismatches 82; Indels 29;
                                                         Score 301.5; DB 9;
Pred. No. 1.7e-20;
1; Mismatches 82;
                                                         Indels 29;
                                                                                             Length
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RESULT 13 US-10-176-921-262

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US-10-176-918-262
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US-10-176-918-262
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Best Local S
Matches 78
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SEQ ID NO 262
LENGTH: 242
TYPE: PRT
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Publication No. US20030027275A1
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CURRENT FILING DATE: 2002-06-20
Prior Application removed - See File Wrapper or
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                                                                                                          121 -SWSQAIQSWFDEYNDFDFGVGFKTPNAVVGHYTQVVWYSSYLVGCGNAYCFN-QKVLKY 178
                                        179 YYVCQYCPAGNWANRLYVPYEQGAPCASC -- PDNCDDGLC 216
                                                                                 108 FTPRHAITAWYNETQFYDF--DSLSCSRVCGHYTQLVWANSFYVGCAVAMCPNLGGASTA 165
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les 78; Conserv
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IFVCNYGPAGNFAN--MPPYARGESCSLCSKEEKCVKNLC 203
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Watanabe, Colin K
Wood, William
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Gurney, Austin L.
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                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                           21.0%;
                                                                                                                                                                                                                                                                                                                         Score 301.5; DB 9;
pred. No. 1.7e-20;
1; Mismatches 82;
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                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                    67
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Prior Application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 550

SEQ ID NO 262

LENGTH: 242

TYPE: PRT
ORGANIZM: Homo Sapien
US-10-176-921-262
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                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                      Sequence 262, A Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C28B: US/10/176,921 CURRENT APPLICATION NUMBER: US/10/176,921 CURRENT FILING DATE: 2002-06-20
                                                                                                                                                                                                                                  APPLICANT: Baker, Kevin P. APPLICANT: Beresini, Maur
                                                                                                                                                                                                                                                                                                                                                                                                                                             179 YYVCQYCPAGNWANRLYVPYEQGAPCASC -- PDNCDDGLC 216
                                                                                                                                                                                                                                                                                                                                                                                                        166 IFVCNYGPAGNFAN -- MPPYARGESCSLCSKEEKCVKNLC 203
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        121 -SWSQAIQSWFDEYNDFDFGVGPKTPNAVVGHYTQVVWYSSYLVGCGNAYCPN-QKVLKY 178
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ALKNKFSCLWILGLCLVATTSSKIPS--ITDPHFI------DNCIEAHNEWRGKVN 49
                                                                     Gerritsen, Mary E. Goddard, Audrey Godowski, Paul J. Gurney, Austin L.
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Godowski, Paul J.
Gurney, Austin L.
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Filvaroff, Ellen
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o. US20030027276A1
                                                                                                                                                    Gao, Wei-Qiang
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                                  Smith, Victoria
                                                   Sherwood, Steven
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b. US20030032155A1
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Pred. No. 1.7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 242;
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US-10-137-865-262
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US-10-140-474-262
                                      US-10-140-474-262
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Best Local S
Matches 78
Query Match
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APPLICANT:
APPLICANT:
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                                                        ORGANISM: Homo Sapien
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Watanabe, Colin K
Wood, William
                                                                                                                                                                                                                                                                                                                                                                                     Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
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Stewart, Timothy A.
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21.0%; Score 301.5;
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CURRENT APPLICATION NUMBER: US/10/137,865
CURRENT FILING DATE: 2002-05-03
Prior Application removed - See Palm or File Wrapper
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 262
LENGTH: 242
TYPE: PRT
CURRENT APPLICATION NUMBER: US/10/140,474
CURRENT FILING DATE: 2002-05-06
Prior Application removed - See Palm or File Wrapper
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 262
LENGTH: 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 262, Application US/10140474 Publication No. US20030032156A1 GENERAL INFORMATION:
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APPLICANT: Wood, William
APPLICANT: Zhang, Zemin
APPLICANT: Zhang, Zemin
ATTILE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C154
                                                                                                                                                                                                TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C162
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Length 242;

121 - SWSOAIOSWFDEYNDFDFGVGPKTDNAVVGHYTOVVWYSSYIVGCGNAVCDN-OKVIKY 178	2 ALKNKESCHMILGLCLVATTSSKIPS ITDPHFI	Best Local Similarity 35.5%; Pred. No. 1.7e-20; Matches 78; Conservative 31; Mismatches 82; Indels 29; Gaps 7 8 ALETTAMTLEPVLLFLVAGLLPSFPANEDKDPAFTALLTTQTQVQREIVNKHNELRRAVS 67
121 -SWSQAIQSWFDEYNDFDFGVGPKTPNAVVGHYTQVVWYSSYLVGCGNAYCPN-QKVLKY 178	: RGKVN 49 SSAPS 120 - GGIKS 107	9; Gaps RRAVS 67